

(FILE 'HOME' ENTERED AT 13:20:48 ON 21 MAR 2001)

FILE 'CAPLUS, USPATFULL, WPIDS, DGENE, EUROPATFULL, JAPIO, MEDLINE'  
ENTERED AT 13:21:31 ON 21 MAR 2001

L1	45 S MAE (W) II
L2	1 S L1 AND DPD
L3	2350 S DPD
L4	104 S L3 AND (MUTATION OR POLYMORPHISM)
L5	44 S L4 NOT PY>1996
L6	0 S 1-10 IBIB ABS

=> d his

(FILE 'HOME' ENTERED AT 14:07:01 ON 16 MAR 2001)

FILE 'CAPLUS, USPATFULL, WPIDS, DGENE, EUROPATFULL, JAPIO' ENTERED AT 14:07:34 ON 16 MAR 2001

L1 353 S DIHYDROPYRIMIDINE (W) DEHYDROGENASE  
L2 288659 S MUTATION OR POLYMORPHISM  
L3 71 S L2 AND L1  
L4 1 S L3 AND MAEIII  
L5 0 S L1 AND ALTERNATE (W) SPLIC?  
L6 41 S L1 AND SCREEN?

=> s l6 not py>1996

3 FILES SEARCHED...

L7 9 L6 NOT PY>1996

=> d ibib abs 1-

YOU HAVE REQUESTED DATA FROM 9 ANSWERS - CONTINUE? Y/(N):y

L7 ANSWER 1 OF 9 CAPLUS COPYRIGHT 2001 ACS

ACCESSION NUMBER: 1996:690425 CAPLUS

DOCUMENT NUMBER: 125:325384

TITLE: A point mutation in an invariant splice donor site leads to exon skipping in two unrelated Dutch patients with **dihydropyrimidine dehydrogenase** deficiency

AUTHOR(S): Vreken, P.; Van Kuilenburg, B. P.; Meinsma, R.; Smit, G. P. A.; Bakker, H. D.; De Abreu, R. A.; van Gennip, A. H.

CORPORATE SOURCE: Acad. Med. Cent., Univ. Amsterdam, Amsterdam, 1100 DE, Neth.

SOURCE: J. Inherited Metab. Dis. (1996), 19(5), 645-654  
CODEN: JIMDDP; ISSN: 0141-8955

DOCUMENT TYPE: Journal

LANGUAGE: English

AB **Dihydropyrimidine dehydrogenase** (DPD) deficiency is an autosomal recessive disease characterized by thymine-uraciluria and assocd. with a variable clin. phenotype. To identify the mol. defect underlying complete DPD deficiency in a Dutch patient previously shown to have a 165 base pair deletion in the mature DPD mRNA, the authors cloned the genomic region encompassing the skipped exon and its flanking intron sequences. Sequence anal. revealed that the patient was homozygous for a single G.fwdarw. A point mutation in the invariant GT dinucleotide splice donor site downstream of the skipped exon. The same mutation was identified in another, unrelated, Dutch patient. Because this mutation destroys a unique MaeII restriction site, rapid **screening** using restriction enzyme cleavage of the amplified genomic region encompassing this mutation is possible. Anal. of 50 controls revealed no individuals heterozygous for this mutation.

L7 ANSWER 2 OF 9 CAPLUS COPYRIGHT 2001 ACS

ACCESSION NUMBER: 1996:483095 CAPLUS

DOCUMENT NUMBER: 125:139772

TITLE: Molecular basis of the human **dihydropyrimidine dehydrogenase** deficiency and 5-fluorouracil toxicity

AUTHOR(S): Wei, Xiaoxiong; McLeod, Howard L.; McMurrough, Julieann; Gonzalez, Frank J.; Fernandez-Salguero, Pedro

CORPORATE SOURCE: Laboratory of Molecular Carcinogenesis, National Institutes of Health, Bethesda, MD, 20892, USA

SOURCE: J. Clin. Invest. (1996), 98(3), 610-615

CODEN: JCINAO; ISSN: 0021-9738

DOCUMENT TYPE: Journal

LANGUAGE: English

AB **Dihydropyrimidine dehydrogenase** (DPD) deficiency constitutes an inborn error in pyrimidine metab. assocd. with thymine-uraciluria in pediatric patients and an increased risk of toxicity in cancer patients receiving 5-fluorouracil (5-FU) treatment. The mol.